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RAW SEQUENCE LISTING

PATENT APPLICATION: US/09/902,525

DATE: 04/26/2002

TIME: 14:28:57

Input Set : N:\Crf3\RULE60\09902525.raw  
Output Set: N:\CRF3\04262002\I902525.raw

SEQUENCE LISTING

1 (1) GENERAL INFORMATION:

2 (i) APPLICANT: Recombinant Biocatalysis, Inc.  
3 (ii) TITLE OF INVENTION: THERMOSTABLE PHOSPHATASES  
4 (iii) NUMBER OF SEQUENCES: 54  
5 (iv) CORRESPONDENCE ADDRESS:  
6 (A) ADDRESSEE: Fish & Richardson P.C.  
7 (B) STREET: 4225 Executive Square, Suite 1400  
8 (C) CITY: La Jolla  
9 (D) STATE: CA  
10 (E) COUNTRY: USA  
11 (F) ZIP: 92037  
12 (v) COMPUTER READABLE FORM:  
13 (A) MEDIUM TYPE: Diskette  
14 (B) COMPUTER: IBM Compatible  
15 (C) OPERATING SYSTEM: Windows 95  
16 (D) SOFTWARE: FastSEQ for Windows Version 2.0b  
17 (vi) CURRENT APPLICATION DATA:

C--> 18 (A) APPLICATION NUMBER: US/09/902,525  
C--> 19 (B) FILING DATE: 09-Jul-2001

20 (C) CLASSIFICATION:

21 (vii) PRIOR APPLICATION DATA:  
22 (A) APPLICATION NUMBER: US/09/202,681A  
23 (B) FILING DATE: 23-Dec-1999  
25 (A) APPLICATION NUMBER: 08/666,857  
26 (B) FILING DATE: 19-JUN-1996  
27 (A) APPLICATION NUMBER: 60/033,752  
28 (B) FILING DATE: 19-JUN-1997  
29 (viii) ATTORNEY/AGENT INFORMATION:  
30 (A) NAME: Haile, Ph.D., Lisa A.  
31 (B) REGISTRATION NUMBER: 38,347  
32 (C) REFERENCE/DOCKET NUMBER: 09010/015W01

33 (ix) TELECOMMUNICATION INFORMATION:

34 (A) TELEPHONE: 619/678-5070  
35 (B) TELEFAX: 619/678-5099  
36 (C) TELEX:

37 (2) INFORMATION FOR SEQ ID NO: 1:

38 (i) SEQUENCE CHARACTERISTICS:  
39 (A) LENGTH: 52 base pairs  
40 (B) TYPE: nucleic acid  
41 (C) STRANDEDNESS: single  
42 (D) TOPOLOGY: linear  
43 (ii) MOLECULE TYPE: cDNA

ENTERED

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Input Set : N:\Crf3\RULE60\09902525.raw  
 Output Set: N:\CRF3\04262002\I902525.raw

44 (xi) SEQUENCE DESCRIPTION: SEQ ID NO: 1:  
 45 CCGAGAATTCAATTAAAGAGG AGAAATTAAC TATGGGGCA GGTCCGAAAA GG 52  
 47 (2) INFORMATION FOR SEQ ID NO: 2:  
 48 (i) SEQUENCE CHARACTERISTICS:  
 49 (A) LENGTH: 31 base pairs  
 50 (B) TYPE: nucleic acid  
 51 (C) STRANDEDNESS: single  
 52 (D) TOPOLOGY: linear  
 53 (ii) MOLECULE TYPE: cDNA  
 54 (xi) SEQUENCE DESCRIPTION: SEQ ID NO: 2:  
 55 CCGAGGATCC TCACCGCCCC CTGCGGGTGC G 31  
 57 (2) INFORMATION FOR SEQ ID NO: 3:  
 58 (i) SEQUENCE CHARACTERISTICS:  
 59 (A) LENGTH: 52 base pairs  
 60 (B) TYPE: nucleic acid  
 61 (C) STRANDEDNESS: single  
 62 (D) TOPOLOGY: linear  
 63 (ii) MOLECULE TYPE: cDNA  
 64 (xi) SEQUENCE DESCRIPTION: SEQ ID NO: 3:  
 65 CCGAGAATTCAATTAAAGAGG AGAAATTAAC TATGTTGGAT ATACTGCTTG TT 52  
 67 (2) INFORMATION FOR SEQ ID NO: 4:  
 68 (i) SEQUENCE CHARACTERISTICS:  
 69 (A) LENGTH: 32 base pairs  
 70 (B) TYPE: nucleic acid  
 71 (C) STRANDEDNESS: single  
 72 (D) TOPOLOGY: linear  
 73 (ii) MOLECULE TYPE: cDNA  
 74 (xi) SEQUENCE DESCRIPTION: SEQ ID NO: 4:  
 75 CCGACGATCC TTATTTTTA ACCAAATGTT CC 32  
 77 (2) INFORMATION FOR SEQ ID NO: 5:  
 78 (i) SEQUENCE CHARACTERISTICS:  
 79 (A) LENGTH: 52 base pairs  
 80 (B) TYPE: nucleic acid  
 81 (C) STRANDEDNESS: single  
 82 (D) TOPOLOGY: linear  
 83 (ii) MOLECULE TYPE: cDNA  
 84 (xi) SEQUENCE DESCRIPTION: SEQ ID NO: 5:  
 85 CCGACAATTCAATTAAAGAGG AGAAATTAAC TATGATGATG GAATTCACTC GC 52  
 87 (2) INFORMATION FOR SEQ ID NO: 6:  
 88 (i) SEQUENCE CHARACTERISTICS:  
 89 (A) LENGTH: 32 base pairs  
 90 (B) TYPE: nucleic acid  
 91 (C) STRANDEDNESS: single  
 92 (D) TOPOLOGY: linear  
 93 (ii) MOLECULE TYPE: cDNA  
 94 (xi) SEQUENCE DESCRIPTION: SEQ ID NO: 6:  
 95 CCGAGGATCC CTACAGTCT AAAAGTCTTT TA 32  
 97 (2) INFORMATION FOR SEQ ID NO: 7:  
 98 (i) SEQUENCE CHARACTERISTICS:

## RAW SEQUENCE LISTING

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Input Set : N:\Crf3\RULE60\09902525.raw  
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99                   (A) LENGTH: 52 base pairs  
 100                  (B) TYPE: nucleic acid  
 101                  (C) STRANDEDNESS: single  
 102                  (D) TOPOLOGY: linear  
 103                (ii) MOLECULE TYPE: cDNA  
 104               (xi) SEQUENCE DESCRIPTION: SEQ ID NO: 7:  
 105                CCGACAATTG ATTAAAGAGG AGAAATTAAC TATGAGAACCTTGACAATAA AC       52  
 107 (2) INFORMATION FOR SEQ ID NO: 8:  
 108               (i) SEQUENCE CHARACTERISTICS:  
 109                (A) LENGTH: 31 base pairs  
 110                (B) TYPE: nucleic acid  
 111                (C) STRANDEDNESS: single  
 112                (D) TOPOLOGY: linear  
 113               (ii) MOLECULE TYPE: cDNA  
 114               (xi) SEQUENCE DESCRIPTION: SEQ ID NO: 8:  
 115                CCGAGGATCC TTACACCCAC AGAACCCCTTA C                           31  
 117 (2) INFORMATION FOR SEQ ID NO: 9:  
 118               (i) SEQUENCE CHARACTERISTICS:  
 119                (A) LENGTH: 52 base pairs  
 120                (B) TYPE: nucleic acid  
 121                (C) STRANDEDNESS: single  
 122                (D) TOPOLOGY: linear  
 123               (ii) MOLECULE TYPE: cDNA  
 124               (xi) SEQUENCE DESCRIPTION: SEQ ID NO: 9:  
 125                CCGAGAATTG ATTAAAGAGG AGAAATTAAC TATGAAAGGA AAGTCTCTTG TT       52  
 127 (2) INFORMATION FOR SEQ ID NO: 10:  
 128               (i) SEQUENCE CHARACTERISTICS:  
 129                (A) LENGTH: 31 base pairs  
 130                (B) TYPE: nucleic acid  
 131                (C) STRANDEDNESS: single  
 132                (D) TOPOLOGY: linear  
 133               (ii) MOLECULE TYPE: cDNA  
 134               (xi) SEQUENCE DESCRIPTION: SEQ ID NO: 10:  
 135                CCGAGGATCC TCAAGCTTCC TGGAGAATCA A                           31  
 137 (2) INFORMATION FOR SEQ ID NO: 11:  
 138               (i) SEQUENCE CHARACTERISTICS:  
 139                (A) LENGTH: 52 base pairs  
 140                (B) TYPE: nucleic acid  
 141                (C) STRANDEDNESS: single  
 142                (D) TOPOLOGY: linear  
 143               (ii) MOLECULE TYPE: cDNA  
 144               (xi) SEQUENCE DESCRIPTION: SEQ ID NO: 11:  
 145                CCGAGAATTG ATTAAAGAGG AGAAATTAAC TATGCCAAGA AATATGCCG CT       52  
 147 (2) INFORMATION FOR SEQ ID NO: 12:  
 148               (i) SEQUENCE CHARACTERISTICS:  
 149                (A) LENGTH: 34 base pairs  
 150                (B) TYPE: nucleic acid  
 151                (C) STRANDEDNESS: single  
 152                (D) TOPOLOGY: linear

RAW SEQUENCE LISTING  
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153 (ii) MOLECULE TYPE: cDNA  
154 (xi) SEQUENCE DESCRIPTION: SEQ ID NO: 12:  
155 CCGAGGATCC TTAAGGCTTC TCGAGGTGGG GGTT 34  
157 (2) INFORMATION FOR SEQ ID NO: 13:  
158 (i) SEQUENCE CHARACTERISTICS:  
159 (A) LENGTH: 54 base pairs  
160 (B) TYPE: nucleic acid  
161 (C) STRANDEDNESS: single  
162 (D) TOPOLOGY: linear  
163 (ii) MOLECULE TYPE: cDNA  
164 (xi) SEQUENCE DESCRIPTION: SEQ ID NO: 13:  
165 CCGACAATTG ATTAAAGAGG AGAAATTAAC TATGTATAAA TGGATTATTG AGGG 54  
167 (2) INFORMATION FOR SEQ ID NO: 14:  
168 (i) SEQUENCE CHARACTERISTICS:  
169 (A) LENGTH: 31 base pairs  
170 (B) TYPE: nucleic acid  
171 (C) STRANDEDNESS: single  
172 (D) TOPOLOGY: linear  
173 (ii) MOLECULE TYPE: cDNA  
174 (xi) SEQUENCE DESCRIPTION: SEQ ID NO: 14:  
175 CCGAGGACTA AACATAGTCT AAGTAATTAG C 31  
177 (2) INFORMATION FOR SEQ ID NO: 15:  
178 (i) SEQUENCE CHARACTERISTICS:  
179 (A) LENGTH: 52 base pairs  
180 (B) TYPE: nucleic acid  
181 (C) STRANDEDNESS: single  
182 (D) TOPOLOGY: linear  
183 (ii) MOLECULE TYPE: cDNA  
184 (xi) SEQUENCE DESCRIPTION: SEQ ID NO: 15:  
185 CCGAGAATTTC ATTAAAGAGG AGAAATTAAC TATGAGAATC CTCCTCACCA AC 52  
187 (2) INFORMATION FOR SEQ ID NO: 16:  
188 (i) SEQUENCE CHARACTERISTICS:  
189 (A) LENGTH: 31 base pairs  
190 (B) TYPE: nucleic acid  
191 (C) STRANDEDNESS: single  
192 (D) TOPOLOGY: linear  
193 (ii) MOLECULE TYPE: cDNA  
194 (xi) SEQUENCE DESCRIPTION: SEQ ID NO: 16:  
195 CCGAGGATCC TCACAGGCTC AGAACGCCTTT G 31  
197 (2) INFORMATION FOR SEQ ID NO: 17:  
198 (i) SEQUENCE CHARACTERISTICS:  
199 (A) LENGTH: 54 base pairs  
200 (B) TYPE: nucleic acid  
201 (C) STRANDEDNESS: single  
202 (D) TOPOLOGY: linear  
203 (ii) MOLECULE TYPE: Genomic DNA  
204 (xi) SEQUENCE DESCRIPTION: SEQ ID NO: 17:  
205 CCGAGAATTTC ATTAAAGAGG AGAAATTAAC TATGGAAAAC TTAAAAAAGT ACCT 54  
207 (2) INFORMATION FOR SEQ ID NO: 18:

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 PATENT APPLICATION: US/09/902,525 TIME: 14:28:57

Input Set : N:\Crf3\RULE60\09902525.raw  
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208 (i) SEQUENCE CHARACTERISTICS:  
 209 (A) LENGTH: 31 base pairs  
 210 (B) TYPE: nucleic acid  
 211 (C) STRANDEDNESS: single  
 212 (D) TOPOLOGY: linear  
 213 (ii) MOLECULE TYPE: cDNA  
 214 (xi) SEQUENCE DESCRIPTION: SEQ ID NO: 18:  
 215 CCGAGGATCC TCACCGCCCC CTGCGGGTGC G 31  
 217 (2) INFORMATION FOR SEQ ID NO: 19:  
 218 (i) SEQUENCE CHARACTERISTICS:  
 219 (A) LENGTH: 783 base pairs  
 220 (B) TYPE: nucleic acid  
 221 (C) STRANDEDNESS: single  
 222 (D) TOPOLOGY: linear  
 223 (ii) MOLECULE TYPE: Genomic DNA  
 224 (ix) FEATURE:  
 225 (A) NAME/KEY: Coding Sequence  
 226 (B) LOCATION: 1...780  
 227 (xi) SEQUENCE DESCRIPTION: SEQ ID NO: 19:  
 228 ATG AGG GGG AGC GGA GTG CGG ATA CTT CTC ACC AAC GAT GAC GGC ATC 48  
 229 Met Arg Gly Ser Gly Val Arg Ile Leu Leu Thr Asn Asp Asp Gly Ile  
 230 1 5 10 15  
 231 TTT GCC GAG GGT CTG GGG GCT CTG CGC AAG ATG CTG GAG CCC GTG GCT 96  
 232 Phe Ala Glu Gly Leu Gly Ala Leu Arg Lys Met Leu Glu Pro Val Ala  
 233 20 25 30  
 234 ACC CTT TAC GTG GTG GCT CCG GAC CGA GAG CGT AGC GCG GCC AGC CAT 144  
 235 Thr Leu Tyr Val Val Ala Pro Asp Arg Glu Arg Ser Ala Ala Ser His  
 236 35 40 45  
 237 GCT ATC ACC GTT CAC CGC CCC CTG CGG GTG CGG GAG GCG GGT TTT CGC 192  
 238 Ala Ile Thr Val His Arg Pro Leu Arg Val Arg Glu Ala Gly Phe Arg  
 239 50 55 60  
 240 AGC CCC AGG CTT AAA GGC TGG GTA GTG GAC GGT ACC CCG GCC GAC TGC 240  
 241 Ser Pro Arg Leu Lys Gly Trp Val Val Asp Gly Thr Pro Ala Asp Cys  
 242 65 70 75 80  
 243 GTC AAG CTG GGC CTG GAG GTA CTT TTG CCC GAA CGT CCA GAT TTC CTG 288  
 244 Val Lys Leu Gly Leu Glu Val Leu Leu Pro Glu Arg Pro Asp Phe Leu  
 245 85 90 95  
 246 GTT TCG GGC ATA AAC TAC GGG CCC AAC CTG GGT ACC GAC GTA CTT TAC 336  
 247 Val Ser Gly Ile Asn Tyr Gly Pro Asn Leu Gly Thr Asp Val Leu Tyr  
 248 100 105 110  
 249 TCC GGC ACC GTC TCG GCG GCC ATA GAA GGG GTA ATT AAC GGC ATT CCC 384  
 250 Ser Gly Thr Val Ser Ala Ala Ile Glu Gly Val Ile Asn Gly Ile Pro  
 251 115 120 125  
 252 TCG GTG GCC GTA TCT TTG GCC ACG CGG CGG GAG CCG GAC TAT ACC TGG 432  
 253 Ser Val Ala Val Ser Leu Ala Thr Arg Arg Glu Pro Asp Tyr Thr Trp  
 254 130 135 140  
 255 GCG GCC CGG TTC GTC CTG GTC CTG CTG GAG GAA CTG CGA AAA CAC CAA 480  
 256 Ala Ala Arg Phe Val Leu Val Leu Glu Glu Leu Arg Lys His Gln  
 257 145 150 155 160

**VERIFICATION SUMMARY**

PATENT APPLICATION: US/09/902,525

DATE: 04/26/2002

TIME: 14:28:58

Input Set : N:\Crf3\RULE60\09902525.raw  
Output Set: N:\CRF3\04262002\I902525.raw

L:18 M:220 C: Keyword misspelled or invalid format, [(A) APPLICATION NUMBER:]  
L:19 M:220 C: Keyword misspelled or invalid format, [(B) FILING DATE:]